

Neural Networks as a tool for parameter estimation in mega-pixel data sets

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ABSTRACT

We present a neural net algorithm for parameter estimation in the context of large cosmological data sets. Cosmological data sets present a particular challenge to pattern-recognition algorithms since the input patterns (galaxy redshift surveys, maps of cosmic microwave background anisotropy) are not fixed templates overlaid with random noise, but rather are random realizations whose information content lies in the correlations between data points. We train a “committee” of neural nets to distinguish between Monte Carlo simulations at fixed parameter values. Sampling the trained networks using additional Monte Carlo simulations generated at intermediate parameter values allows accurate interpolation to parameter values for which the networks were never trained. The Monte Carlo samples automatically provide the probability distributions and truth tables required for either a frequentist or Bayesian analysis of the one observable sky. We demonstrate that neural networks provide unbiased parameter estimation with comparable precision as maximum-likelihood algorithms but significant computational savings. In the context of CMB anisotropies, the computational cost for parameter estimation via neural networks scales as $N^{3/2}$. The results are insensitive to the noise levels and sampling schemes typical of large cosmological data sets and provide a desirable tool for the new generation of large, complex data sets.

Subject headings: methods: data analysis — (cosmology:) cosmic microwave background — (cosmology:) cosmological parameters

1. Introduction

A fundamental question in cosmology is the origin and evolution of large scale structure in the universe. The standard model for this evolution is the gravitational growth and collapse of initially small perturbations in the primordial density distribution. This picture is supported by the detection of primordial Cosmic Microwave Background (CMB) temperature anisotropies at a level of approximately one part in 10^5 by the Cosmic Background Explorer satellite and a series of ground-based and balloon-borne experiments. Small perturbations on the matter and energy density in the early universe are reflected in the temperature distribution of the CMB, providing a “snapshot” of conditions the early universe while the perturbations were still in the linear regime.

One angular scale of particular interest is the horizon size at the surface of last scattering, the epoch when the universe cooled sufficiently to form neutral hydrogen and allow the CMB photons to propagate freely. Causally-connected regions at the surface of last scattering, as viewed from the present epoch, subtend an angle

$$\theta \sim 1^\circ 7 \Omega_0^{1/2} \left(\frac{1100}{1 + z_{ls}} \right)^{1/2}$$

where z_{ls} is the redshift at last scattering and Ω_0 is the total density of the universe relative to the critical (closure) density. Anisotropy on scales larger than $\sim 2^\circ$ reflect perturbations larger than the particle horizon and thus probe the primordial density distribution. On scales smaller than 2° , causal mechanisms become important and modify the primordial density in model-specific ways.

Oscillations in the coupled photon-baryon fluid within the primordial potential wells prior to decoupling lead to signature oscillations (“acoustic peaks”) in the angular power spectrum of the CMB (Peebles and Yu 1970). A precise determination of the power spectrum in this regime can probe a wealth of information about the early universe. For instance, the angular scale of the first acoustic peak depends primarily on the angle subtended by the Jeans mass at last scattering (Hu and Sugiyama 1995). It is thus a direct probe of the large-scale geometry of the universe and hence the density parameter Ω_0 . The width of the acoustic peak depends on the sound speed at decoupling, and in turn on the baryon density and Hubble constant as $\Omega_b h^2$. A measurement of this quantity could be compared with similar determinations based on primordial nucleosynthesis (Copi *et. al.* 1995; Olive *et. al.* 2000). See *e.g.* (Hu 1999), for a review of how the different physical processes in the early universe leave their imprint on the CMB anisotropies. Measurements of the CMB anisotropy on medium angular scales thus offer an elegant determination of such parameters as the curvature, density, matter density, baryon density, Hubble constant, cosmological constant, density perturbation index n , scalar/tensor ratio, and ionization history of the universe.

From the one observable sky, we want to infer the values of these cosmological parameters with minimal uncertainty in the shortest possible time. Theoretical models do not predict a specific template for the CMB anisotropy (a hot spot at this location, a cold spot over there), but rather predict a statistical distribution usually expressed in terms of the angular power spectrum. Deriving the power spectrum from the data (or more generally, deriving model parameters directly from the sky maps) involves accounting for angular correlations between pixels, precluding use of simple linear least-squares techniques. The accepted standard in the CMB community has been the generalization of least-squares techniques as implemented in maximum likelihood algorithms (see, e.g., (Górski et al. 1996; Tegmark, Taylor, & Heavens 1997; Bond, Jaffe, & Knox 1998; Borrill 1999)).

The simplest method of parameter estimation uses a goodness-of-fit test to compare a set of observables $y_i \pm \sigma_i$ measured at a set of positions x_i to a theoretical model Γ_i . If we have N_d data points y_i and N_p parameters p_j , we define

$$\chi^2 = \sum_{i=1}^{N_d} \left(\frac{y_i - \Gamma_i}{\sigma_i} \right)^2, \quad (1)$$

where

$$\Gamma_i(x) = \sum_{j=1}^{N_p} p_j X_j(x_i) \quad (2)$$

is function of the parameters p and some fixed basis functions $X(x)$. We obtain the “best-fit” parameter values by minimizing χ^2 with respect to the parameters,

$$\frac{\partial \chi^2}{\partial p_j} = 0 \quad (3)$$

for the j^{th} parameter p_j . The least-squares system in Eq. 3 has the solution

$$p_j = \sum_{k=1}^{N_p} (\mathbf{A}^{-1})_{jk} B_k \quad (4)$$

where

$$\mathbf{A}_{jk} = \sum_{i=1}^{N_d} \frac{X_j(x_i) X_k(x_i)}{\sigma_i^2} \quad (5)$$

is an $N_p \times N_p$ matrix, and

$$B_k = \sum_{i=1}^{N_d} \frac{y_i X_k(x_i)}{\sigma_i^2} \quad (6)$$

is a vector of length N_p .

If the data points are not independent, this relatively simple calculation becomes much more costly. Covariance between the observed data points can result from instrumental artifacts (correlated noise, instrumental resolution, oversampled data) or from correlations in the underlying signal (for instance, measuring in real space a signal whose components are independent in Fourier space). Equation 1 can be generalized to include the effects of covariance,

$$\chi^2 = \sum_{i=1}^{N_d} \sum_{j=1}^{N_d} (y_i - \Gamma_i) (\mathbf{M}^{-1})_{ij} (y_j - \Gamma_j), \quad (7)$$

where

$$\mathbf{M}_{ij} = \langle (y_i - \langle \Gamma_i \rangle) (y_j - \langle \Gamma_j \rangle) \rangle \quad (8)$$

is the $N_d \times N_d$ covariance matrix and the brackets denote an ensemble average.

Conjugate gradient techniques can solve for χ^2 without expliciting solving for \mathbf{M}^{-1} and thus avoiding the $O(N_d^3)$ operations this would incur. But if the covariance matrix \mathbf{M} depends on the parameters p_j , then minimizing χ^2 will produce biased estimates for p_j . Maximum-likelihood parameter estimation provide a tool to overcome this limitation. For a multivariate Gaussian distribution, the probability of obtaining the observed data y_i given a set of parameters p_j is

$$\mathcal{L} = P(y|p) = (2\pi)^{-N_d/2} \frac{\exp(-\frac{1}{2}\chi^2)}{|\mathbf{M}|^{1/2}} \quad (9)$$

where χ^2 is defined in Eq. 7. The “best” choice of parameters is that which maximizes the likelihood function \mathcal{L} . The curvature of the likelihood surface about the maximum defines the uncertainty in the fitted parameters,

$$\delta p_j \geq \sqrt{(\mathbf{F}^{-1})_{jj}} \quad (10)$$

where

$$\mathbf{F}_{ij} = \langle \frac{\partial^2 L}{\partial p_i \partial p_j} \rangle \quad (11)$$

is the Fisher information matrix (Kendall & Stuart 1969) and $L = -\log(\mathcal{L})$ (see Bunn & Sugiyama 1995; Vogeley & Szalay 1996; Tegmark et al. 1997; Bond et al. 1998).

The maximum likelihood estimator is unbiased and asymptotically approaches the equality in Eq. 10. However, these advantages come at a steep price: both the χ^2 and the determinant calculation in Eq. 9 scale as $O(N_d^3)$, making brute-force techniques computationally infeasible. For $N_d > 10^6$ the time required is measured in years, even on the most powerful supercomputers. A number of authors have suggested ways around this problem. Karhunen-Loève eigenvalue techniques produce moderate data compression, reducing the N_d original

data points to $N' \approx N_d/10$ eigenmodes (Bond 1994; Bunn & Sugiyama 1995; Tegmark et al. 1997). However, estimating cosmological parameters from the smaller set of eigenmodes still scales as $(N')^3$ operations, making such techniques undesirable for mega-pixel data sets.

Oh et al. (1998) derive a method for likelihood evaluation using a Newton-Raphson quadratic iteration scheme. This method utilizes a conjugate gradient algorithm to evaluate χ^2 . The determinant is first approximated using azimuthal symmetry of the noise matrix (appropriate for full-sky CMB maps), then corrected using Monte Carlo simulations. The method provides a nearly minimum-variance estimate of the angular power spectrum for CMB anisotropy maps in $O(N_d^2)$ operations and $O(N_d^{3/2})$ storage; cosmological parameters can then be derived by comparing the power spectrum to various models. Although this algorithm is fast enough for mega-pixel data sets, it has several weaknesses. Since it is a Newton-Raphson iterative scheme, it requires a sufficiently good starting estimate to guarantee convergence to true maximum. It is also optimized to estimate the power spectrum, rather than the underlying cosmological parameters – when used as a root-finding technique in parameter space, the radius of convergence is small and the problems associated with parameter covariance become severe.

The last few years have seen the development of a number of techniques designed to overcome the N_d^3 problem, most of them focusing on estimating the power spectrum. These techniques include Szapudi *et. al.* (2000), who have developed a method based on using the two-point correlation function. Doré *et. al.* (2001) relies on an hierarchical implementation of the usual quadratic estimator to the power spectrum. Both methods are known to scale as N_d^2 , with the first hoping to be improved to $N_d \log N_d$ while the second may scale as N_d (with a large prefactor). There is also the Monte Carlo estimator to the power spectrum developed by Hivon *et. al.* 2001, which has been used (Netterfield *et. al.* 2001) to analysis the BOOMERANG data (deBernadis *et. al.* 2000). All these approximate techniques rely on first estimating the power spectrum and then using this estimated power spectrum to determine the most likely cosmological parameters. Douspis *et. al.* (2001) have shown that the data sets are not fully described by just their band power sets.

Borrill (1998) offers a global solution to bound the likelihood. This method uses Gaussian quadrature to bound the likelihood at *any* point in parameter space (not just near the likelihood maximum); it is thus well suited to search parameter space using the minimum-variance direct pixel basis. However, the method requires $O(N_d^{7/3})$ operations for each likelihood evaluation and is thus significantly slower than the method of Oh et al. (1998). More importantly, it can only provide bounds on the likelihood, fixing $\log(\mathcal{L})$ to accuracy of a few percent. Since $\log(\mathcal{L}) > N_d$ (a large number), errors of a few percent can create significant bias in the location of the likelihood maximum.

The Microwave Anisotropy Probe (MAP), launched in the summer of 2001, will measure the full sky in 5 frequency bands with over a million pixels per band (Bennett *et. al.* 1995). The Planck Surveyor mission, scheduled later in the decade, will produce maps with over 10^7 pixels (Bersanelli *et. al.* 1996). Maps of such size can not be analyzed with brute-force maximum likelihood techniques.

Figure 1 shows typical information flow for cosmological parameter estimation. We compare data (CMB sky maps) to a parameterized model using some mathematical “machinery” to derive a set of parameters describing the data. Although least squares methods and their generalization to maximum-likelihood techniques are common choices for this machinery, they are not the only choices possible. For mega-pixel data sets, these deterministic methods are computationally infeasible. Neural net algorithms provide an alternative machinery for parameter estimation in large, complex data sets.

Neural networks have been used previously in astronomy for galaxy classification (Lahav *et. al.* 1996; Andreon *et. al.* 2000) and periodicity analysis of unevenly sampled data as applied to stellar light curves (Tagliaferri *et. al.* 2000). They have also been used to analyze stellar spectra (Bailer-Jones *et. al.* 1997; Bailer-Jones *et. al.* 1998; Bailer-Jones 2000), with results comparable to traditional methods. However, Bailer-Jones *et. al.* compared data to a deterministic model (stellar spectra), whereas cosmological applications examine random patterns drawn from parameterized stochastic models. We demonstrate the generality of our algorithm by considering different problems with the same network architecture.

We use neural networks as a complementary approach to cosmological parameter estimation in large data sets. They are a forward algorithm in that they “learn” the differences between two different parameter data points by being trained on sets of simulated data sets drawn at the specified parameter points. We find the computational cost for training the network, in the context of CMB anisotropy, requires $O(N^{3/2})$ operations and thus provides a substantial improvement over brute-force maximum-likelihood methods.

Neural networks do not require that we specify one single statistic of *a priori* interest. As the network is trained, it determines how it will discriminate. The information required to separate different parameter points comes from the training set simulations. After selecting a pair of parameter points, a committee of networks are trained on simulations taken at each of the parameter points. We interpolate by sampling the trained networks with simulated data drawn at parameter points between the training sets. This takes advantage of the fact that for large input patterns, the number of sampling sets will be much less than the number of training passes. The algorithm naturally provides the distributions necessary to understand the confidence levels of the parameter fit when the physical data is analyzed. This sort of sampling of trained networks becomes a key component in utilizing a Bayesian approach

to parameter estimation, see (Christensen and Meyer 2000; Rocha 2000) for discussions of the Bayesian approach in the context CMB anisotropies and (MacKay 1995), along with (Bishop 1995), for neural networks.

We present a brief review of the neural network algorithm and discuss how this model is used for parameter estimation. To demonstrate the feasibility of this algorithm, we show how we can estimate parameters in three different examples. The same neural network and parameter estimation algorithm is used in all the examples. Using these examples, we vary the size and noise characteristics of the simulated data sets to determine the computational scaling.

2. Neural Network Algorithms

Parameter estimation can be thought of as an exercise in pattern recognition, for which an extensive literature exists (see, e.g., McCulloch and Pitts 1943; Rosenblatt 1962; Watanabe 1969; Hopfield 1982; Rumelhart, Hinton, & McClelland 1986; Grossberg 1988). Neural networks, an implementation of parallel distributed processing architecture, are well suited to pattern recognition. Although not widely recognized as such, neural networks are also well suited for the problem of parameter estimation in large, complex data sets.

The basic building block of a neural network is the neuron, consisting of any number of inputs and a single output. A neuron sums its inputs to determine its output via some (non-linear) transfer function. A neural network consists of layers of neurons connected together via matrices of weights (Figure 2). A network is trained to produce a desired output by repeatedly presenting the network with a set of known input patterns, then adjusting the weights until the network produces the desired output for each known input. When the network is later presented with an unknown input, the output will reflect which of the known inputs the unknown input most closely resembles.

Figure 2 shows a typical network architecture. The first layer in a network is its input layer, whose outputs reflect the pattern presented to the network. Any number of hidden layers lie between the input and output layers, from which the results of the network are read. We use one hidden layer and a single output unit. We thus have one matrix of weights connecting the input pattern to the inputs of the hidden layer and another matrix connecting the outputs of the hidden layer to the inputs of the output unit.

The information content of a neural network lies in the matrices of weights connecting layers (analogous to the information content of the covariance matrix for maximum likelihood algorithms). The weights are set during the training process, in which we present the network

with a series of known inputs and adjust the weights to obtain the desired outputs. For a network with N_d input units, we let \mathbf{X}_{patt} , a N_d element vector, represent an input pattern. The output of the network can be viewed as a function of the input pattern: $o = o(\mathbf{X}_{\text{patt}})$. For a given pattern we associate an output target t_{patt} (in our case a single value, but for a more general network, this would be a N_{output} vector). A training algorithm is a method for adjusting the weights to minimize the total error $E = \sum_{\text{patt}} E_{\text{patt}}$, where

$$E_{\text{patt}} = \frac{1}{2} (o(\mathbf{X}_{\text{patt}}) - t_{\text{patt}})^2 \quad (12)$$

We use the back propagation method (Rumelhart, D.E., Hinton, G.E., and Williams, R.J. 1986). According to this method, we present a series of N_{Train} patterns to the network, one at a time. For each pattern, we compare the output with the associated target and determine the error. We use this error to adjust the weights connecting the hidden layer to the output layer, and then back-propagate it to the weights between the hidden and input layers. Note that the training depends on the difference between the desired and actual outputs - no *a priori* statistical test for the inputs is specified.

Neural nets achieve computational savings by training with simulated data sets, as opposed to inverting a large matrix. The computational cost to train a neural net is found to be

$$N_{\text{CPU}} \propto N_d^\alpha, \quad (13)$$

where the exponent α depends on the particular problem. There are two main contributions to this cost: the array multiplication needed to sum the weights to evaluate a single pattern, times the number of training patterns required. In general we find $1 \leq \alpha < 3$, depending on the specific problem. This is never worse than standard maximum likelihood methods, and at best the computational cost grows linearly with the input pattern size. For the specific case of parameter estimation in mega-pixel CMB maps, the computational cost scales as $N_{\text{CPU}} \propto N_d^{1.5}$. Neural networks provide significant computational savings over both standard maximum-likelihood algorithms and the N^2 scaling for the fastest known approximate method.

3. Neural Networks and Parameter Estimation

Neural networks can be trained to estimate the value of a continuous parameter, and can reliably interpolate to parameter values intermediate between the training values. Though this idea is not new to astrophysics, *e.g.* (Bailer-Jones *et. al.* 1997; Bailer-Jones *et. al.* 1998; Bailer-Jones 2000), our method is fundamentally different from that of Bailer-Jones *et. al.*

in that our patterns are random samples drawn from a parameterized parent population. The Bailer-Jones *et. al.* method is a matter of template matching; randomness only enters their input patterns as instrument noise. For CMB and other cosmological data, the patterns themselves are intrinsically random. Nonetheless, using the same basic neural network architecture, we can train the networks to discriminate stochastic patterns that differ according to the parent population from which they are drawn.

We start by training a network to differentiate between simulated data sets (including instrument noise and other artifacts) generated at a pair of discrete parameter values. The back-propagation adjusts the weights until the network outputs target value 0 when presented with the first set of patterns, and target value 1 when presented with the second. In practice, since the information distinguishing different parameter values is in the correlations, not the actual pixel values, any single network will not train to a sufficient degree. We improve the situation by training a small committee of networks and polling them to get a consensus opinion. Now we find simulations generated at the training parameter values produce two well defined peaks. Simulations generated with an intermediate parameter value (never present in training data) yield outputs peaking somewhere in between, depending on whether the new parameter value is closer to the first or second training value (see Fig 3).

We quantify this behavior by presenting the trained network with a set of new inputs drawn from a grid of intermediate parameter values, and derive for each intermediate parameter value the corresponding probability distribution of output values. When the networks are later presented with an unknown pattern, each distribution gives the probability that the unknown pattern was generated with a parameter value corresponding to that grid point. The interpolated parameter is the probability-weighted mean. The grid samples all use the same trained networks; the sampling of the networks at different grid points is faster than the training since we usually need many more training sets than sampling sets, thus no great computational cost is incurred. Although we focus below on estimating a single parameter, the method is readily extended to multi-parameter fits.

The simulations are viewed as random variables $\mathbf{X}(p)$ taken from an underlying parameterized probability distribution, with p the parameter. Assuming the data lies between two extreme values $p^{(0)}$ and $p^{(1)}$ we train a network to the target $t^{(0)} = 0$ for realizations $\mathbf{X}(p^{(0)})$ and $t^{(1)} = 1$ for $\mathbf{X}(p^{(1)})$ by repeatedly presenting the network with new samples at $p^{(0)}$ and $p^{(1)}$ and back propagating the error. Once trained, the network will give an output $o(\mathbf{X})$ between 0 and 1 for any input parameter value. If $p < p^{(0)}$, the output clips at 0, while if $p > p^{(1)}$ the output clips at 1.

Once we have trained a network, we can present additional, statistically independent samples drawn at $p^{(0)}$ and $p^{(1)}$. Figure 3a shows the output distributions for 1000 patterns

of each parameter. We see the network has successfully trained in that the $p^{(0)}$ distribution is peaked at $o = 0$ while the $p^{(1)}$ samples at $o = 1$. To be able to interpolate to intermediate values, we will need to present samples drawn at intermediate parameter values. Fig 3b shows the output distributions for an additional 1000 samples each for two parameters, one just a little larger than $p^{(0)}$ and one a little smaller than $p^{(1)}$. These distribution also show the same tendency to peak close to the limits of 0 and 1, but not as strongly as those drawn at the trained parameters. In effect, the network is choosing which of the training parameters these new patterns, for which it was never trained, most closely resemble. At it stands now, this tendency makes it hard to construct the probability distributions we need for parameter estimation.

By using a committee of networks, we take advantage of this peaking tendency. We want to determine the committee consensus and from this get the distributions we seek. The first step is converting the continuous output value into a discrete *truth values* 0 or 1. For each trained network, we associate a midpoint value o_{mid} and for any input pattern \mathbf{X} , we define its truth value according to

$$\mathbf{t} \equiv \mathbf{t}(\mathbf{X}) = \begin{cases} 0; & o(\mathbf{X}) \leq o_{\text{mid}} \\ 1; & o(\mathbf{X}) > o_{\text{mid}} \end{cases} . \quad (14)$$

We interpret $\mathbf{t}(\mathbf{X}) = 0$ as indicating the pattern was drawn from the parent population with parameter $p^{(0)}$, and similarly we associate $\mathbf{t} = 1$ with $p^{(1)}$.

To determine o_{mid} , we present N samples drawn at $p^{(0)}$ and at $p^{(1)}$. For each of these sets and any \tilde{o}_{mid} , we obtain the truth values $\mathbf{t}_i^{(0)}$ and $\mathbf{t}_i^{(1)}$, $i = 1, \dots, N$. With this, $\mathbf{n}^{(0)} = \sum_i (1 - \mathbf{t}_i^{(0)})$ is the number patterns drawn at $p^{(0)}$ correctly identified as drawn at $p^{(0)}$ and $\mathbf{n}^{(1)} = \sum_i \mathbf{t}_i^{(1)}$ similarly at $p^{(1)}$. We chose o_{mid} to maximize $f_C = \frac{1}{2N}(\mathbf{n}^{(0)} + \mathbf{n}^{(1)})$ and refer to f_C as the *fraction correct*, our main measure of how well a network has trained. In Fig 3a, o_{mid} is marked with the vertical line and we find $f_C = 94\%$.

We note f_C allows us to determine the optimal number of training passes, N_{Train} . Starting with an initially randomized network, as the network trains, we intermittently pause the training and sample the network to determine f_C . It steadily increases to a maximum value and then levels out: the minimum for the training error E has been reached. We take N_{Train} to be just where this plateau starts.

For each network, any given input pattern is converted into discrete truth values \mathbf{t} . We now form a committee of such networks, where the only difference between the networks is the initial randomization of the weights. We find committee sizes $N_{\text{net}} \sim 50$ sufficient. After presenting any given pattern \mathbf{X} to the committee, we have the collection of truth values $\mathbf{t}(\mathbf{X})_m$, $m = 1, \dots, N_{\text{net}}$. We view each \mathbf{t}_m as the vote from network m as to whether

the pattern resembled those drawn at $p^{(0)}$ or $p^{(1)}$. The committee consensus is formed by generating the *average truth value*

$$\bar{\mathbf{t}}(\mathbf{X}) = \frac{1}{N_{\text{net}}} \sum_{m=1}^{N_{\text{net}}} \mathbf{t}(\mathbf{X})_m. \quad (15)$$

Figure 3c shows the distribution of average truth values for the same set of samples drawn at $p^{(0)}$ and $p^{(1)}$ used in Fig 3a. They are now even more sharply peaked about $\bar{\mathbf{t}} = 0$ and $\bar{\mathbf{t}} = 1$ and in terms of the average truth value, $f_C = 100\%$. More important are the distributions displayed in Fig 3d. These are for the same intermediate samples as used in Fig 3b; we have now two well defined distributions with peaks intermediate to the peaks for the $p^{(0)}$ and $p^{(1)}$ samples. This is the general trend when we work in terms of the average truth value: as the parameter p is swept from $p^{(0)}$ to $p^{(1)}$, we get well defined distributions whose peak moves from $\bar{\mathbf{t}} \sim 0$ to $\bar{\mathbf{t}} \sim 1$.

We utilize this behavior to have our networks interpolate parameter values for which it was *never* trained. We need a function that maps an average truth value to an estimated parameter: $\bar{\mathbf{t}} \mapsto \mathbf{p}(\bar{\mathbf{t}})$. We built this function by sampling the committee of networks with samples drawn at parameter values intermediate to the training values (computationally a small cost compared with training a network). These in turn are used to determine the optimal function $\mathbf{p}(\bar{\mathbf{t}})$. The results of such sampling also automatically provide the distributions needed for either a frequentist's or Bayesian analysis. Selecting K parameter values uniformly distributed between the training values, $p^k = p^{(0)}, p^{(0)} + \Delta p, p^{(0)} + 2\Delta p, \dots, p^{(1)}$, we generate N samples at each of these parameter values. The samples are presented to the committee of networks and thus for each sample \mathbf{X}_i drawn at each parameter value p^k , its average truth value $\bar{\mathbf{t}}_i^k$ is computed. We determine the parameter estimation function $\mathbf{p}(\bar{\mathbf{t}})$ by minimizing the distances between estimated and true parameter values, *i.e.* the error

$$E_{\mathbf{p}} = \frac{1}{2} \sum_k \sum_i (\mathbf{p}(\bar{\mathbf{t}}_i^k) - p^k)^2. \quad (16)$$

The $\bar{\mathbf{t}}$'s will take on discrete values, $\bar{\mathbf{t}} = 0, 1/N_{\text{net}}, 2/N_{\text{net}}, \dots, 1$, and thus so will the parameter estimation function: $\mathbf{p}(\bar{\mathbf{t}}) = \mathbf{p}_j$ when $\bar{\mathbf{t}} = j/N_{\text{net}}$. Let n_j^k be the number of samples drawn at p^k with $\bar{\mathbf{t}}_i^k = j/N_{\text{net}}$. Then we can re-write our error as $E_{\mathbf{p}} = \frac{1}{2} \sum_{k,j} n_j^k (\mathbf{p}_j - p^k)^2$. Minimizing this yields

$$\frac{\partial E_{\mathbf{p}}}{\partial \mathbf{p}_j} = 0 \quad \Rightarrow \quad \mathbf{p}_j = \frac{\sum_k n_j^k p^k}{\sum_k n_j^k}, \quad (17)$$

i.e., each binned value of the parameter estimation function is the parameter weighted average of the number of patterns of each sample that fall into that bin. Figures 3c and 3d are examples the n_j^k distributions.

Given an observed data set, represented as the input pattern \mathbf{X}_{obs} , we present it to the committee of networks and obtain its average truth value $\bar{\mathbf{t}}_{\text{obs}}$. From this, we get the estimated parameter value \mathbf{p}_{obs} . We may now generate a set of samples at this parameter value, $\{\mathbf{X}_i(\mathbf{p}_{\text{obs}})\}$, and estimate their parameters, $\mathbf{p}_{i,\text{fit}}$. The distribution of the $\mathbf{p}_{i,\text{fit}}$ ’s provides a self-consistent method for converging to the final estimation of the parameter. We begin with a wide separation between training values in an attempt to bracket the unknown input. We know the initial range is too narrow if either $\mathbf{p}_{\text{obs}} \sim p^{(0)}$ or $\sim p^{(1)}$, that is, if the output clips at either end of its trained range. The distribution of the $\mathbf{p}_{i,\text{fit}}$ ’s also tells us when we have converged on final answer. We compare the width in the fitted parameter distribution to the separation between training values, and select new (closer) values. We iterate this refinement of $p^{(0)}$ and $p^{(1)}$ until the mean $\bar{\mathbf{p}}_{\text{fit}}$ doesn’t change or the width of the fitted distribution matches the separation of training parameters. In practice convergence typically requires only two or three iterations.

We illustrate how we determine the confidence levels of our parameter estimation via a Bayesian analysis of the results. This is not the only way to proceed; if one preferred, a frequentist’s approach could be taken, similar to the above paragraph. From the Bayesian viewpoint, we are interested in posterior distribution $\mathcal{P}(p|\mathbf{p}_{\text{obs}})$: given the observed/estimated parameter \mathbf{p}_{obs} , what is the probability the true parameter is p . With this, we view the uncertainty in the estimated parameter as

$$\sigma^2(\mathbf{p}_{\text{obs}}) = \int p^2 \mathcal{P}(p|\mathbf{p}_{\text{obs}}) dp - \left(\int p \mathcal{P}(p|\mathbf{p}_{\text{obs}}) dp \right)^2, \quad (18)$$

the width of true parameter values around p_{obs} that have a reasonable probability of being identified as p_{obs} . We can also use this determine the confidence levels. We use Bayes Theorem to express this in terms of the prior distributions:

$$\mathcal{P}(p|\mathbf{p}_{\text{obs}}) = \mathcal{P}(\mathbf{p}_{\text{obs}}|p) \frac{\mathcal{P}(p)}{\mathcal{P}(\mathbf{p}_{\text{obs}})}. \quad (19)$$

All these priors are readily available in the analysis we have developed so far. If there are the same number of patterns in each of the K sample sets, then we have a uniform prior on the parameters p^k we sampled: $\mathcal{P}(p^k) = 1/K$. The probability of getting any particular \mathbf{p}_{obs} is proportional to the total number of $\bar{\mathbf{t}}$ ’s that have the corresponding value and hence $\mathcal{P}(\mathbf{p}_{\text{obs}}) = \frac{1}{KN} \sum_k n_{\text{obs}}^k$. Given the true input parameter value is p^k , the probability that it is identified as \mathbf{p}_{obs} is proportional to the number of samples with an average truth value that corresponds to \mathbf{p}_{obs} : $\mathcal{P}(\mathbf{p}_{\text{obs}}|p^k) = n_{j=\text{obs}}^k/N$. Thus, the probability distribution, Eqn. 19, is $\mathcal{P}(p^k|\mathbf{p}_{\text{obs}}) = n_{\text{obs}}^k / \sum_k n_{\text{obs}}^k$. The resulting expression for the mean true parameter is the same as Eqn. 17, our optimal choice for defining the parameter estimation function. The

68% confidence width for our parameter estimation becomes

$$\sigma^2(\mathbf{p}_{\text{obs}}) = \frac{\sum_k (p^k)^2 n_{\text{obs}}^k}{\sum_k n_{\text{obs}}^k} - (\mathbf{p}_{\text{obs}})^2 \quad (20)$$

It is important to note all the information used to determine this uncertainty in the fit was already generated during the determination of $\mathbf{p}(\bar{\mathbf{t}})$.

We can use this Bayesian analysis to supplement our convergence criterion outlined above. The posterior probability $\mathcal{P}(p|\mathbf{p}_{\text{obs}})$ is zero for $p < p^{(0)}$ or $p > p^{(1)}$, since there are no samples drawn outside the training range. Thus the uncertainty will be artificially suppressed close to these limits. If \mathbf{p}_{obs} is within $\sigma(\mathbf{p}_{\text{obs}})$ of either $p^{(0)}$ or $p^{(1)}$, we need to select new training limits, *i.e.*, we have encountered the clipping previously discussed. If we are clear of either training limit and $\sigma(\mathbf{p}_{\text{obs}})$ is much less than the width of the training interval, we should choose new, closer, training limits.

Whether we chose to do a frequentist’s or Bayesian analysis, this method of using neural networks for parameter estimation benefits from not requiring the *a priori* definition of a statistic or goodness-of-fit function. We only need to be able to simulate the model, often a computational inexpensive task. Neural networks are ideally suited to working with models where the crucial information is in the phase information, *e.g.* topology of the Universe or tests for non-Gaussianity, along with being a faster algorithm for conventional parameter estimation problems.

4. Examples

To illustrate this algorithm, we analyze three examples. The first is the estimation of the frequency of an irregularly sampled noisy sine wave. This serves as a straightforward example of our method and shows how the neural network can recover a parameter value at which it was never trained. Next, we fit the spectral index for CMB anisotropies based on a Gaussian model, an example of the type of problem that will need accurate methods with low computational cost. The final example is parameter estimation in the context of large scale structure surveys, another arena with large data sets. This example illustrates the strength of the neural network to not need an *a priori* statistic.

4.1. Irregularly Sampled Noisy Sine Wave

A common problem in astrophysics is the estimation of periodic signals in noisy, irregularly sampled data sets. Observing limits often produce data cutouts in time series; the

resulting aliasing of power then precludes simple Fourier transforms. Although other techniques such as periodograms exist for this problem, time series provide a simple introduction to the capabilities of neural nets. Our first example uses a time series consisting of $N_d = 256$ points,

$$X_i(\omega) = \sqrt{2}A_s \sin(2\pi\omega t_i + \phi) + n_i \quad (21)$$

where t_i ranges from 0 to t_f . We restrict data coverage by including 10 uniformly spaced gaps of 10 pixels each, plus an additional 24 randomly-placed gaps (Fig 4a). ϕ is a random phase and n_i is Gaussian random noise of zero mean and unit variance. The factor of $\sqrt{2}$ gives signal-to-noise, S/N , of A_s for each observation (Fig 4b).

This sine wave model is somewhere between a template and a random pattern. If $\phi = \text{const.}$, then this would be a matter of template matching, since but for the noise, the values of each of the pixels would be completely determined for each frequency. This is akin to the classification of stellar spectra (Bailer-Jones *et. al.* 1997; Bailer-Jones *et. al.* 1998; Bailer-Jones 2000). We are allowing ϕ to be a random variable and thus the location the peaks and valleys of the sine wave vary from pattern to pattern. Our model has characteristics of a random pattern: the information about the frequency is now in the correlations of the pixel values.

We train 50 networks using initial parameter range $\omega^{(0)} = 4.0$ and $\omega^{(1)} = 8.0$ to produce an output value of 0 for noisy sine waves with $\omega = 4.0$ and output value 1 for $\omega = 8.0$. We then sample the trained networks at intermediate values of ω to generate nine probability distributions $\mathcal{P}(o; \omega_i)$, $\omega_i = 4.0, 4.5, \dots, 8.0$. Finally, we present the networks with 1000 realizations at frequency $\omega_{\text{in}} = 5.60$. Figure 4c shows the distribution of recovered ω_{obs} from these realizations. The mean value is intermediate between the training inputs, with no output close to the training values of 4.0 or 8.0. The distribution shows most of the estimated parameters lie between $\omega^{(0)} = 4.5$ and $\omega^{(1)} = 6.5$, suggesting we repeat the analysis with these new training values. Fig. 4d shows the result after a second iteration. The peak of the distribution of estimated parameter values has shifted somewhat from the previous iteration. Selecting new training values $\omega^{(0)} = 4.6$ and $\omega^{(1)} = 5.8$ to encompass the bulk of the estimated parameters, we do one more iteration. The peak for this distribution of estimated parameters has not shifted from the previous set of networks and the width of the distribution matches the range of sampled frequencies. The distribution peaks at $\omega = \omega_{\text{in}}$, with fitted mean $\bar{\omega}_{\text{obs}} = 5.56 \pm 0.11$.

Note that we correctly recover ω_{in} despite never training any network at any stage to this value. Neural networks provide an unbiased estimator for a continuous parameter without requiring prior knowledge of the parameter value.

4.2. Microwave Background Anisotropy

Neural networks can also be applied to more challenging computational tasks. Deriving cosmological parameters from maps of the cosmic microwave background usually involves maximum likelihood algorithm whose computational cost (N^3) makes them prohibitive for mega-pixel maps. The same neural net topology used for our time series example provides accurate, rapid parameter estimation for CMB maps as well.

On angular scales $\theta > 2^\circ$, anisotropy in the cosmic microwave background corresponds to primordial density perturbations with scale-free power spectrum $P_k \propto k^n$, where k is the wavenumber and n is the power-law index. We simulate the COBE-DMR full-sky maps of the CMB anisotropy parameterized by the index n (Bond and Efstathiou 1987). We smooth full-sky maps with a Gaussian profile with a FWHM of 7° to include the effect of the radiometer horn profile and add Gaussian noise with an variance of $(20 \text{ mK})^2/N_{\text{obs},i}$ to each pixel to account for the instrument noise (Bennett *et. al.* 1996).

Foreground emission from our Galaxy dominates the COBE data near the Galactic plane, rendering it unusable for cosmological analyses. We use the galaxy cut template of (Banday *et. al.* 1997) to excise pixels with significant Galactic emission. The cut sky represents an additional challenge for standard maximum-likelihood analyses. In the absence of this cut, the data sets represent full-sky coverage and can be decomposed in terms of orthogonal spherical harmonics. The resulting coefficients yield the power spectrum of the CMB and hence the spectral index n . Once the galaxy cut is imposed, the spherical harmonic functions are no longer orthogonal on the remaining pixels. Any attempt to obtain a harmonic expansion will result in the aliasing of power between modes and an inaccurate power spectrum. Though a new orthogonal set of basis functions can be computed for the cut sky (Górski, 1994), this is an N^3 problem as well. Since neural networks estimate cosmological parameters using the real-space pixel values, they need not take the detour through the power spectrum, and do not suffer aliasing of power. We simply impose the cut for galactic emission and train each network using only the remaining high-latitude pixels. As the network is trained, it automatically learns the effect of cuts in the data, without requiring any symmetries in the cut data (see *e.g.* (Oh *et. al.* 1999)).

We generate simulated COBE maps using sky pixels of size $2.5^\circ \times 2.5^\circ$ each. After the galaxy cut, this leaves an input pattern of $N_d = 3881$ pixels. We use $N_{\text{Hidden}} = 600$ and $N_{\text{Train}} = 12000$ (determined to be the optimal choice) and train 50 networks over the parameter range $n^{(0)} = 0.0$ and $n^{(1)} = 2.0$. Figure 5 plots the probability distribution of n_{obs} for a set of 1000 samples of $n_{\text{in}} = 1.40$ (dotted line). This distribution matches with separation of the training sets and we need this only training iteration. We recover $\bar{n}_{\text{obs}} = 1.30$, with 68% of the samples between 1.07 and 1.51. In terms of our Bayesian

analysis, for $n \sim 1$, we determine $\sigma = 0.35$, in agreement with the traditional maximum likelihood analysis of the COBE-DMR 2 year data (Górski *et. al.* 1994).

Note that the neural network algorithm recovered the correct spectral index even though none of the networks used were trained at this value. The uncertainty, derived from the width of the probability distribution of n_{obs} , is comparable to the value predicted by the maximum likelihood method. Neural networks can recover cosmological parameters from CMB data sets with comparable precision as maximum likelihood techniques, but using $N^{1.5}$ calculations instead of N^3 .

4.3. Large-Scale Structure

Large-scale structure provides a third example that combines parameter estimation and phase features. Redshift surveys such as the 2-Degree Field and the Sloan Digital Sky Survey measure the redshift and position on the sky of a large number of galaxies ($N \sim 10^6$), sampling the quasi-linear regime $\sim 100h^{-1}$ Mpc where h is the Hubble constant in units $100 \text{ km s}^{-1} \text{ Mpc}^{-1}$. The observed redshift is the sum of the Hubble flow and the peculiar velocity induced by gravitational acceleration in the evolving density field. Coherent flows on large scales produce artifacts in the redshift distribution compared to real space. Galaxies on the far side of an overdensity tend to flow toward the center (hence toward the observer) so that their peculiar velocities subtract from the Hubble flow, making them appear closer than they really are. Galaxies on the near side move the opposite direction, so their peculiar velocities add to the Hubble flow. The net result is an apparent enhancement in the galaxy density in redshift space on scales of superclusters, compressing the region along the line of sight to the observer. The amplitude of this “bull’s-eye” effect depends on the matter density Ω_m on scales comparable to superclusters of galaxies and can be used to determine Ω_m in model-independent fashion (Paton *et. al.* 1997; Melott *et. al.* 1998).

Estimating Ω_m from distortions in redshift space has several problems in practice. The first is defining a statistic to quantify the bull’s-eye enhancement in concentric rings about the origin. (Melott *et. al.* 1998) use a large number of simulations to develop an empirical statistic defined as the ratio of rms spacing between upcrossings in isodensity contours in the redshift (radial) direction to that in the orthogonal (azimuthal) direction. It is thus a local statistic in that it compares high-density regions only to other nearby regions, and operates only on a single slice of redshift space after smoothing and contouring.

Neural nets, by contrast, offer a *global* test by comparing each region of the density field to all other regions simultaneously, and can easily be extended across the entire three-

dimensional survey. No *a priori* statistic need be identified, nor do neural nets require contouring of the density field, thus avoiding the need to ‘fine-tune’ the selection of contour levels. Figure 6 shows a toy model of the bullseye effect for several samples differing in the amplitude of the radial density enhancements. Our toy model consists of using a Bessel function to modify an otherwise uniform distribution of galaxies, with the amplitude determining the amount of modulation. The inputs to the neural network are density functions, one for each realization. Neural nets, trained to realizations differing only slightly in the amount of modulation, correctly discriminate between them. We do not specify how to tell them apart, but nonetheless, the correct amplitudes are recovered 90% of the time. Note that we test the bullseye effect using the *same* neural nets developed for the noisy sine wave and CMB tests above, simply trained to a different set of reference models. Neural nets are a powerful, versatile tool for a variety of problems in astrophysics.

5. Computational Cost Scaling

Neural nets have many desirable characteristics for parameter estimation with mega-pixel CMB maps. They operate globally on the data and return unbiased estimates of the underlying parameter values. They automatically account for data gaps, instrument noise, and other features peculiar to a particular data set. Most importantly, the computational costs are low enough to allow extension to the mega-pixel data sets expected in the near future.

The dominant contribution to the CPU cost is the array multiplication associated with the weights. This multiplication is performed twice per training pass: once for evaluating the pattern, then again for back propagating the error. Each operation scales as the number of weights. Our total computational cost for training a network is

$$N_{\text{CPU}} = 2 N_{\text{Train}} N_{\text{Hidden}} (N_d + 1) \quad (22)$$

How this cost scales with N_d depends on the problem being considered. We find that $N_{\text{CPU}} \sim N^\alpha$ with $1 < \alpha < 2.5$. For the particular application of mega-pixel CMB maps, $N_{\text{CPU}} \sim N_{\text{Input}}^{1.5}$.

The computational cost depends on how the information content of the input pattern changes as more data points are added. We consider two limiting cases. The first case has fixed signal to noise per data point, so that the signal to noise ratio of the entire data set improves as more points are added. This is analogous to measuring a signal with a noisy detector, and simply adding more observations. The second case keeps the signal to noise ratio of the entire data set fixed, so that the signal to noise ratio per pixel gets worse as

more pixels are added. This is analogous to over-sampling a signal within a limited observing time: as the observations are broken into finer and finer resolution, the observing time per point decreases and the noise per observation gets worse.

We treat these limiting cases using our noisy sine model. For the first case (fixed noise per observation), we simply simulate more data over additional periods of the sine wave, t_f grows with N_d (maintaining both the pattern of regular cutouts and the percentage of random cutouts). The noise per pixel is fixed at $\sigma_i^{\text{noise}} = 1$, independent of N_d . For the second case (fixed noise averaged over the entire data set), we keep the number of periods per data set fixed, $t_f = 1$, but allow the noise amplitude to vary as $\sigma_i^{\text{noise}} = \sigma_0 / \sqrt{N_i^{\text{obs}}}$, and fix σ_0 so $\sigma_i^{\text{noise}} = 1$ when $N_d = 256$.

The number of training passes and the number of hidden units depend on problem considered and the number of input pixels. For each case above, we determine which values give the best N_{CPU} for a fixed training accuracy. For a range of N_{Hidden} , we train multiple networks on a large set of patterns. As they are being trained, we monitor their ability to correctly classify independent samples. Once this ability passes a pre-set threshold, we know N_{Train} for each network, and hence N_{CPU} . We repeat for multiple network to estimate the uncertainty in the CPU cost. (We may also vary other training parameters to assure we have the optimal results.) The results are not strongly dependent on the precise value of N_{Hidden} . Above a minimum value, as N_{Hidden} increases, the number of training passes needed decreases in such a way that N_{CPU} remains constant over a wide range of N_{Hidden} . For these limiting cases, we find $N_{\text{CPU}} \sim N_d$ for fixed noise per pixel, and $N_{\text{CPU}} \sim N_d^{2.5}$ for fixed noise per data set (Figure 7(a)).

The computational costs for CMB analyses lie between these extremes. We derive the scaling for CMB maps by analyzing the CPU costs as progressively larger and larger areas of the sky are covered. In this way, new information is introduced into the data sets as the patch size increases. The S/N ratio per pixel is fixed in this scheme, reflecting the trend in current experiments of scanning ever larger portions of the sky at (roughly) constant S/N ratio per pixel.

We select circular patches of sky centered at the north zenith. The range of patch sizes are chosen to cover 1.5 orders of magnitude in N_d . For each patch size, we proceed as above to determine the optimal CPU cost. The results are shown in Figure 7(b). The computational cost for training a CMB network scales according to $N_{\text{CPU}} \sim N_d^{1.5}$, a considerable improvement over the N_d^3 scaling behavior for a maximum likelihood analysis.

6. Discussion

We have shown neural networks can be used as a tool for astrophysical parameter estimation. For specificity, we have worked in a cosmological context (CMB maps and redshift surveys) where the stochastic nature of the problem is fundamental. The results are insensitive to noise levels and sampling schemes typical of large astrophysical data sets and provide parameter estimation comparable to maximum likelihood techniques. The computational cost is never greater than standard maximum likelihood techniques and in the context for CMB anisotropy maps, we find $N_{\text{CPU}} = O(N_d^{1.5})$.

If we classify parameter estimation techniques as to whether they are forward or reverse algorithms, we see the real strength of neural networks. Maximum-likelihood methods are an example of reverse algorithms. They start with the statistic under consideration and work backwards, inverting a covariance matrix, to the likelihood function used to compare different parameter choices. Forward algorithms provide a way to avoid the high computational costs of inverse methods. Typically, it is much simpler to generate model predictions at each sampled point in parameter space than to compute the matrix inverse and determinant required for maximum likelihood techniques. Forward algorithms trade many realizations of synthetic data sets computed at specific parameter values for the computationally infeasible matrix inversion. Neural networks are such an algorithm; synthetic data sets are used to both train and sample the networks. This gives us our speed improvement.

Since either maximum likelihood or neural networks can be viewed as the “machinery” for parameter estimation, the fundamental information flow stays the same (see figure 1). The statistical confidence levels for the fitted parameters are always accessible. When the “machinery” is sampled with independent synthetic data, we can determine the probabilities for making correct or incorrect parameter identifications. Such sampling also gives us direct access to the statistical power (Phillips and Kogut 2001). While training, the information distinguishing the different parameters is encoded in the weights. Interpreting the resulting weight matrices is not usually possible (as compared to the Fisher matrix, Eqn 11). Using independent sampling of the network to derive the probability distributions needed for, *e.g.* Bayesian analysis, means we do not need direct access to the information in the weight matrices.

A limitation of maximum likelihood methods is their requirement of an *a priori* definition of a goodness-of-fit function. The choice of a goodness-of-fit function is not always obvious and is particularly acute for 2D and 3D surveys. Much of the information lies in the *phase* features of these surveys. Statistical tests can fail badly in detecting phase features, as witness the large literature devoted to the relatively simple problem of edge detection in 2D data sets (see, *e.g.*, (Hough 1962; Davis 1975; Canny 1986)). Topological

tests such as the genus or other Minkowski functionals have been applied to 2- and 3-D maps, (Gott *et. al.* 1990; Kogut *et. al.* 1996) but the relative power of these statistics is poor (Phillips and Kogut 2001). Neural networks, in contrast, do not require specification of a single statistic of *a priori* interest. As the network is trained, it determines how it will discriminate between competing models. The information required to separate different parameter values comes from the training set simulations. Neural networks thus offer a promising approach to cosmological parameter estimation, where the statistical properties of the primordial matter and energy distribution provide one of the few falsifiable tests of the standard inflationary paradigm.

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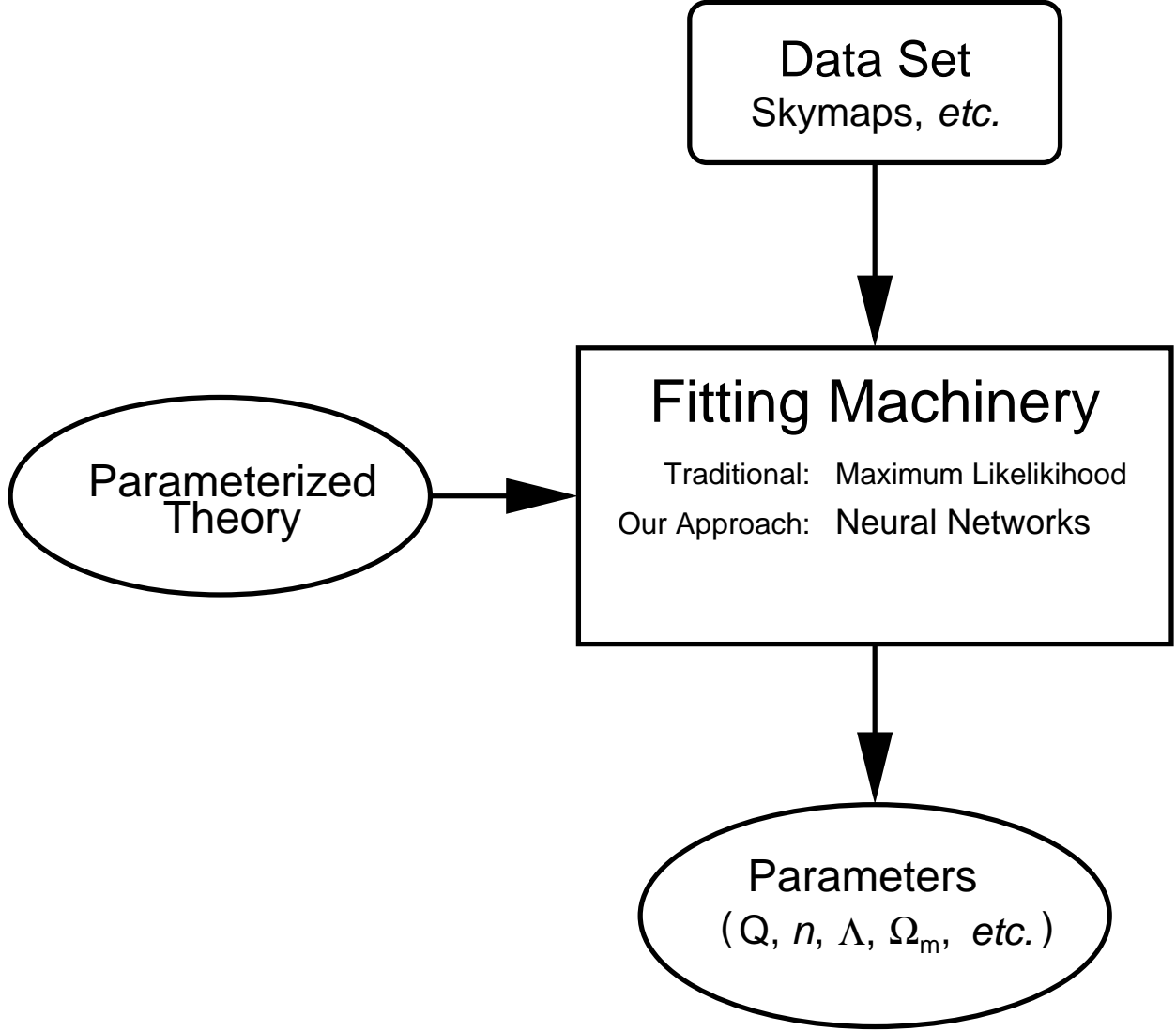


Fig. 1.— Our basic model for parameter estimation from a data set. We have to *a priori* assume a model to compare the observed data set against; what we pay attention to is the machinery for performing this comparison. Maximum likelihood methods, the *de facto* standard in the CMB community must assume a model, just as must be done with currently proposed neural network method.

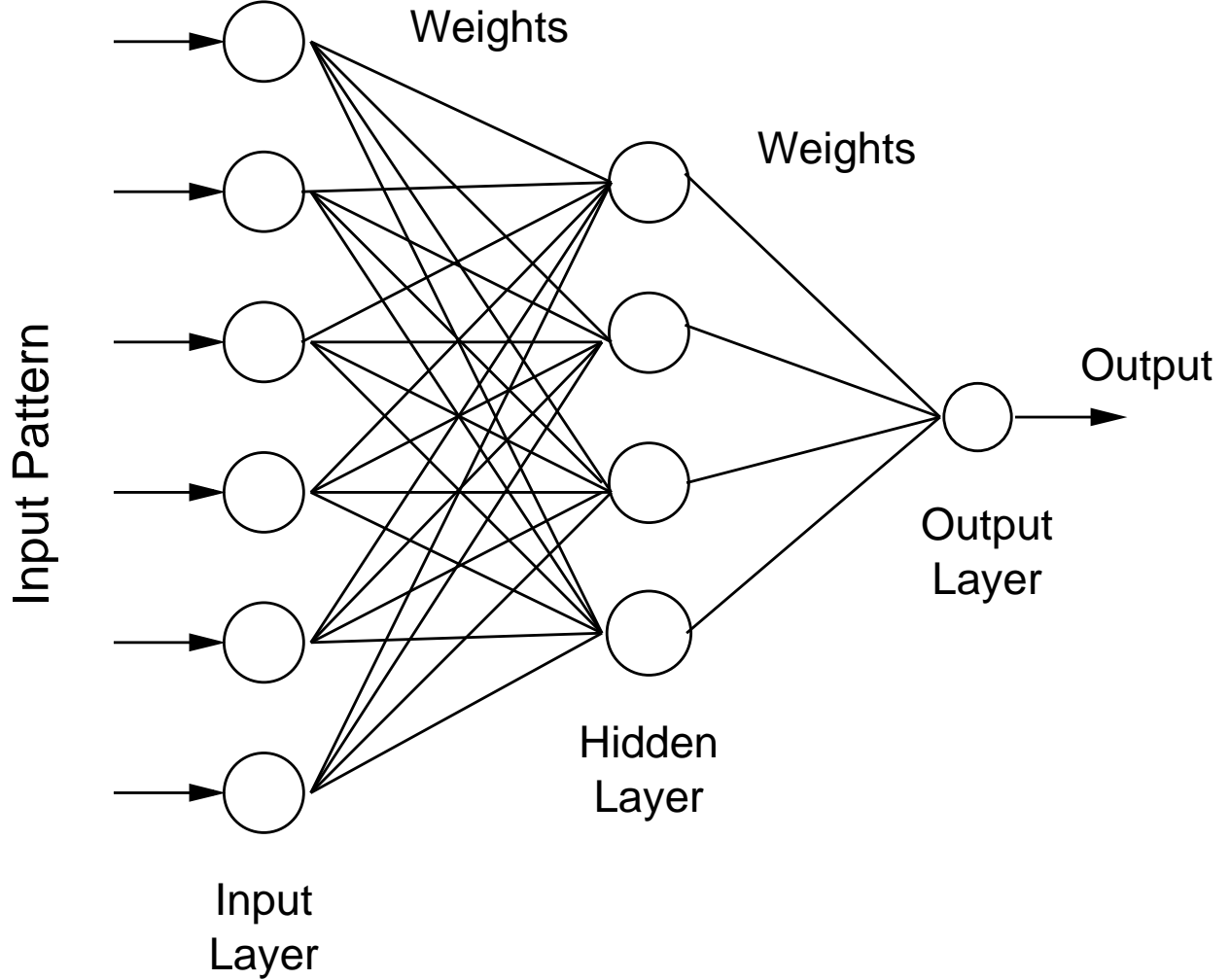


Fig. 2.— Neural network architecture used for parameter estimation. The data (noisy sine wave, CMB maps or galaxy survey) are the input pattern and the neural network feeds forward through the hidden layer to the output unit. The output value is used to derive the probability distributions for the samples and estimated parameter for the observed data set. Typically there are fewer hidden units than input units.

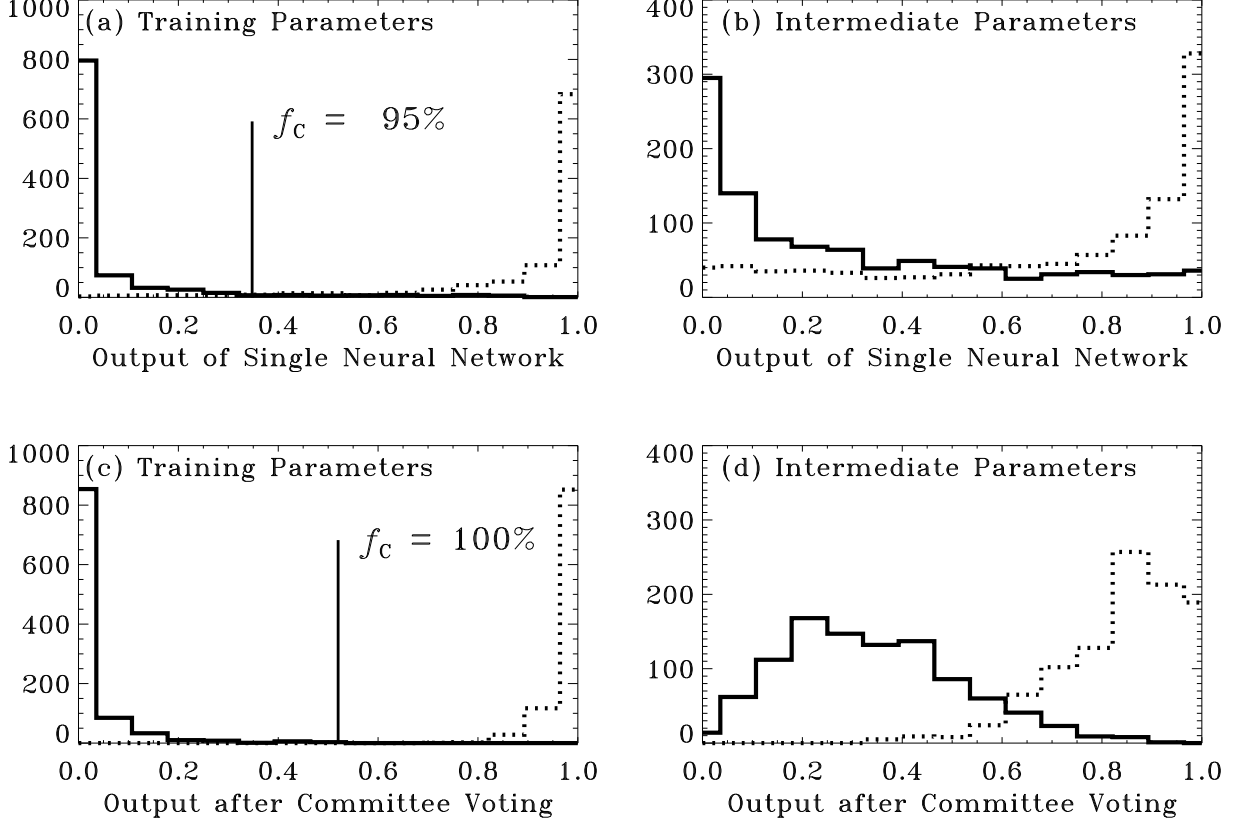


Fig. 3.— Sample neural network output distributions. (a) Solid line is the distribution of output values for an independent set of samples drawn at the same parameter value $p^{(0)}$ for which the network was trained to the target 0. The dotted line is for samples drawn at $p^{(1)}$, the value for the target 1. The vertical line is the midpoint value o_{mid} that maximizes the fraction correct f_c , *i.e.*, all output values $< o_{\text{mid}}$ are identified as being drawn at $p^{(0)}$ while the rest at $p^{(1)}$. (b) The output of the same network, but for samples drawn at parameters intermediate to $p^{(0)}$ and $p^{(1)}$. The solid line is for p close to $p^{(0)}$ and the dotted line for a choice close to $p^{(1)}$. (c) Solid line is the average truth value \bar{t} for the $p^{(0)}$ patterns, averaged over a committee of 50 networks, the only difference between the networks being the initial randomization of the weights. The dotted line is for $p^{(1)}$. (d) The average truth value for the same sets of samples in (b). The averaging has produced two well defined peaks that are cleanly separated. Distributions of \bar{t} like those in (c) and (d) become the basis for predicting which estimated parameters to associate with the average truth values, the output due to presenting a committee of networks with an unknown pattern.

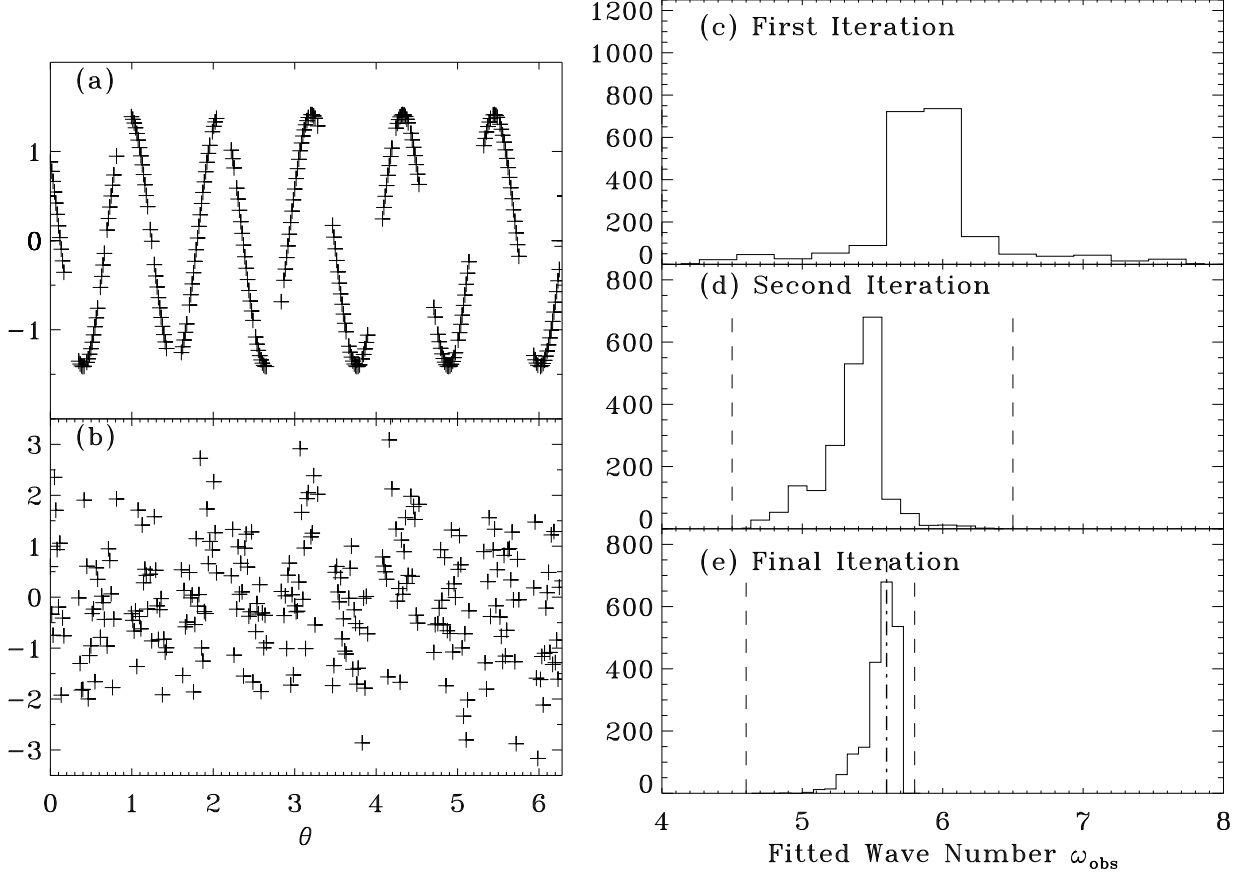


Fig. 4.— Neural net results from noisy time series. (a) Sample sine wave for $\omega_{\text{in}} = 5.60$, including 10 regularly spaced gaps in θ , along with 24 randomly placed gaps. (b) Same sample sine wave, but now including Gaussian noise so that the resulting input network pattern has a $S/N=0.4$ (c) Histogram of fitted ω_{obs} for an initial network training range of $\omega^{(0)} = 4.0$ and $\omega^{(1)} = 8.0$. (d) Fitted values after second iteration. Note that the training values now bracket the distribution returned by the first iteration. Vertical dashed lines show the training frequencies. (e) Fitted values after third (final) iteration. The width of the distribution matches the separation of training values. Note that we correctly recover the input frequency, $\bar{\omega}_{\text{obs}} = 5.56 \pm 0.11$, even though none of the networks were trained at this value.

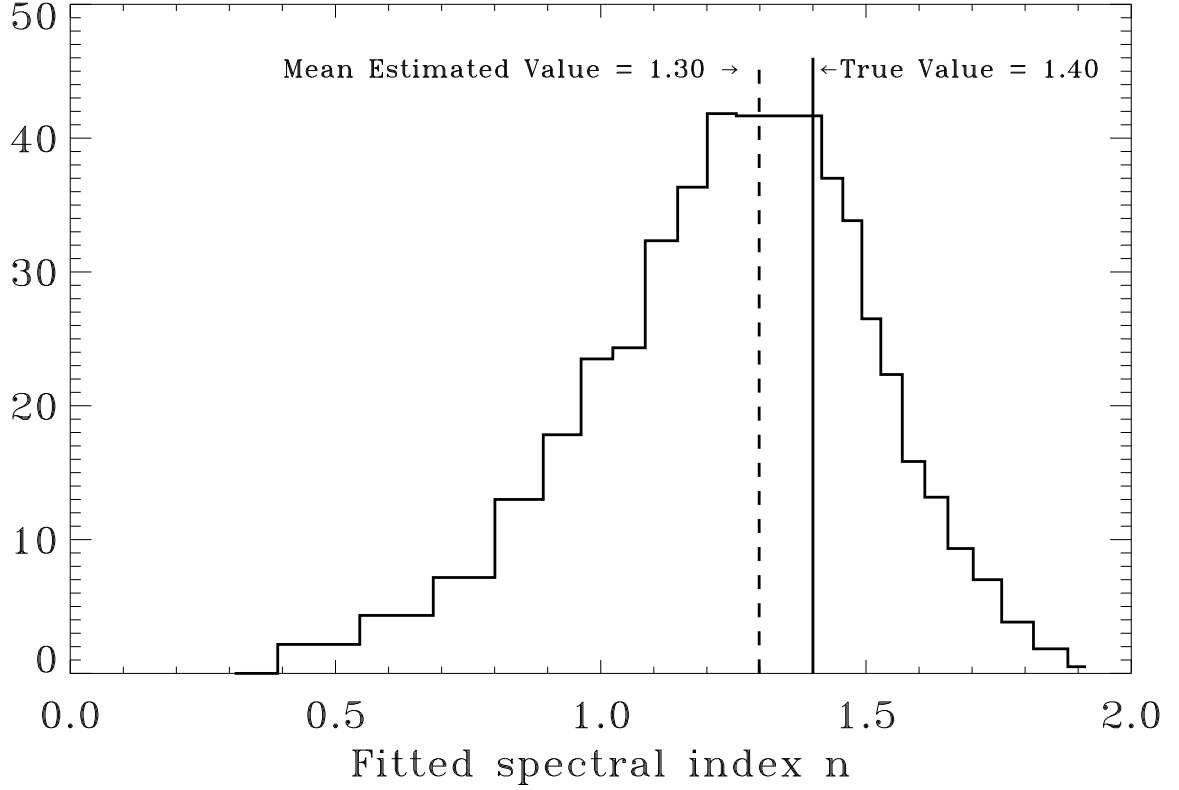


Fig. 5.— Fitted spectral index n_{obs} derived from 1000 realizations of CMB anisotropy sky maps with $n_{\text{in}} = 1.25$. The dotted line is for an initial training range of $n^{(0)} = 0.5$ and $n^{(1)} = 1.5$ while the solid line is the distribution for the final range of $n^{(0)} = 0.8$ and $n^{(1)} = 1.4$. The fitted values correctly peak at the input value (vertical solid line), despite never having trained on this parameter value.

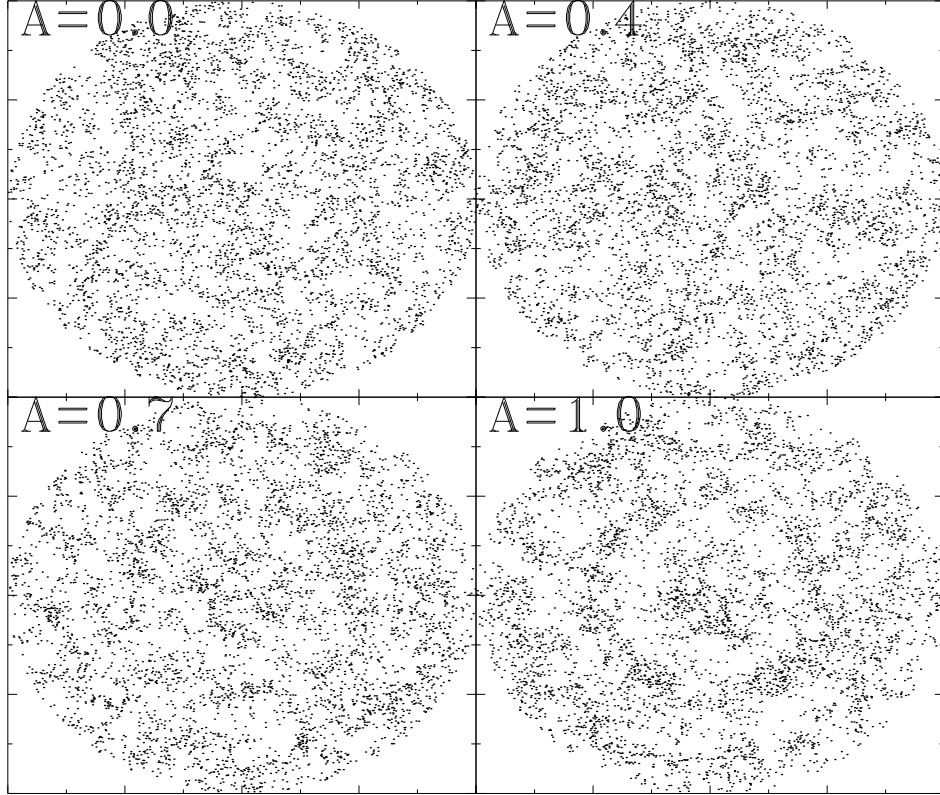


Fig. 6.— Toy models showing changes in the density enhancement A in redshift space due to peculiar velocities (the “bull’s-eye effect”). The same neural nets developed for CMB sky maps successfully discriminate at 90% confidence among all models shown.

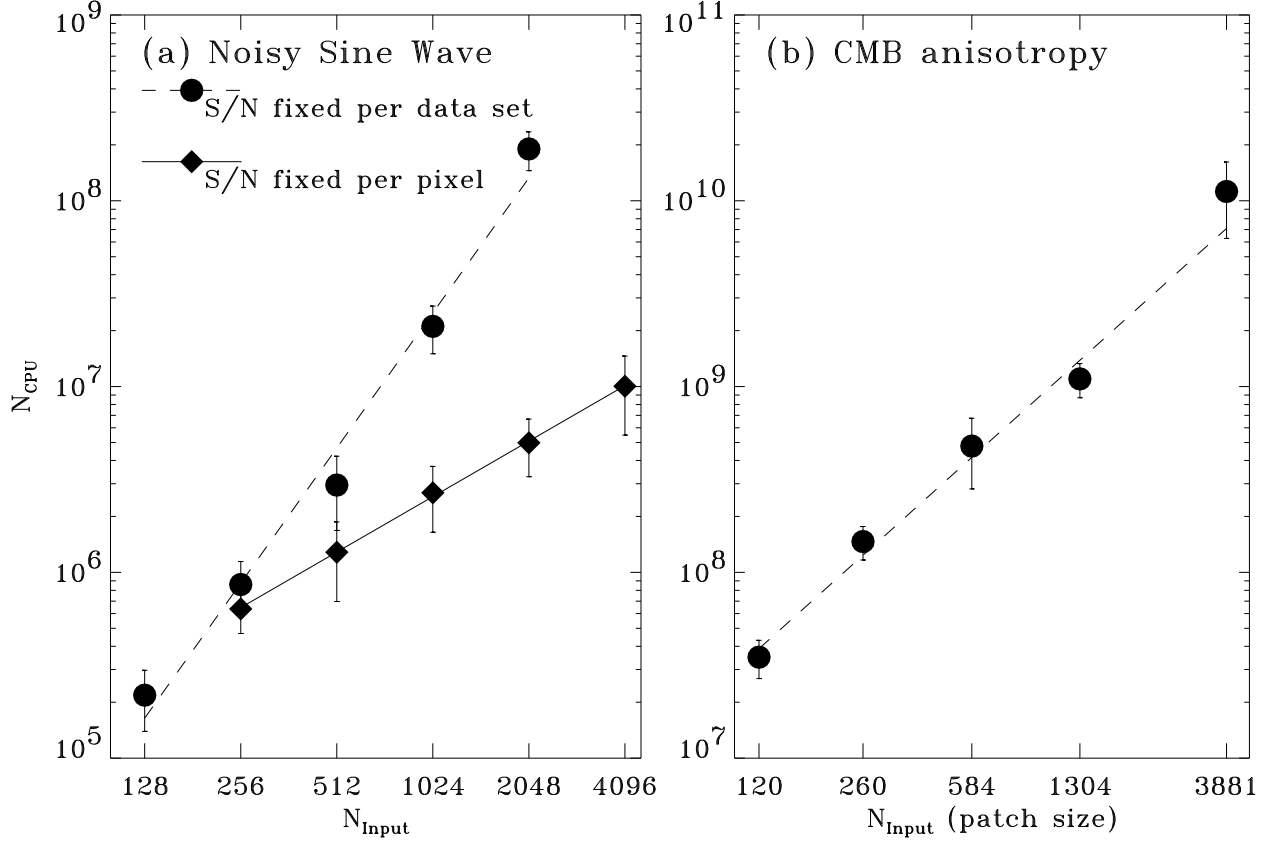


Fig. 7.— CPU cost scaling (a) Scaling of computational costs for the limiting cases of irregularly sampled noisy sine wave. We consider two cases: i) the S/N per pixel is fixed (diamonds/solid line); and ii) the S/N for the entire pattern is fixed and the noise per pixel grows with N_d (circles/dashed line). These limiting cases are well described by power-law fits $N_{\text{CPU}} \sim N_{\text{Input}}^\alpha$ with $\alpha = 1.0$ and 2.5 , respectively. (b) Scaling of computational costs for CMB anisotropy. Working at a fixed sky map resolution, we vary the patch size that is examined. This holds the S/N per pixel fixed, but new information is introduced as the patch size increases. The solid line represents a power law fit of $N_{\text{CPU}} \sim N_d^{1.5}$.